The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

10/698.439A
IFWO
11/2/04

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial	Number: 10/698,439A	CRF Edit Date: ///4/04 Edited by:
	Realigned nucleic acid/amino acid numbers/text text "wrapped" to the next line	in cases where the sequence
	Corrected the SEQ ID NO. Sequence numbers e	dited were:
rene	Inserted or corrected a nucleic number at the end NO's edited:	l of a nucleic line. SEQ ID
/	Deleted:invalid beginning/end-of-file text;_	page numbers
	Inserted mandatory headings/numeric identifiers	, specifically:
	Moved responses to same line as heading/numeric	identifier, specifically:
<u>J</u>	Other: conected spelling of oligonucled conected 21507, 21517 lines	de in Segure 1;



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/698,439A

DATE: 11/04/2004 TIME: 13:02:37

Input Set : A:\PTO.AMC.txt

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4 <110 > APPLICANT: SIGA PHARMACEUTICALS, INC.
      6 <120> TITLE OF INVENTION: New Intergenic and Intragenic Integration Sites for
              Foreign Gene Expression in Recombinant S. Gordonii Strains
      9 <130> FILE REFERENCE: 016921-167
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/698,439A
C--> 11 <141> CURRENT FILING DATE: 2003-11-03
     11 <150> PRIOR APPLICATION NUMBER: PCT/US01/05493
     12 <151> PRIOR FILING DATE: 2001-02-22
     14 <150> PRIOR APPLICATION NUMBER: US 60/184,645
     15 <151> PRIOR FILING DATE: 2000-02-24
     17 <160> NUMBER OF SEQ ID NOS: 49
     19 <170> SOFTWARE: PatentIn version 3.0
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DATE: 11/04/2004 TIME: 13:02:37

PATENT APPLICATION: US/10/698,439A

Input Set : A:\PTO.AMC.txt

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8.8
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PATENT APPLICATION: US/10/698,439A

DATE: 11/04/2004 TIME: 13:02:37

Input Set : A:\PTO.AMC.txt

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PATENT APPLICATION: US/10/698,439A

DATE: 11/04/2004 TIME: 13:02:37

Input Set : A:\PTO.AMC.txt

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DATE: 11/04/2004 PATENT APPLICATION: US/10/698,439A TIME: 13:02:37

Input Set : A:\PTO.AMC.txt

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/698,439A

DATE: 11/04/2004 TIME: 13:02:38

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\11042004\J698439A.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/698,439A

DATE: 11/02/2004 TIME: 13:42:45

Input Set : A:\pto.lm.txt

Output Set: N:\CRF4\11022004\J698439A.raw

- 4 <110> APPLICANT: SIGA PHARMACEUTICALS, INC.
- 6 <120> TITLE OF INVENTION: New Intergenic and Intragenic Integration Sites for
- 7 Foreign Gene Expression in Recombinant S. Gordonii Strains
- 9 <130> FILE REFERENCE: 016921-167
- C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/698,439A
- C--> 12 <141> CURRENT FILING DATE: 2003-11-03
 - 14 <150> PRIOR APPLICATION NUMBER: US 60/184,645
 - 15 <151> PRIOR FILING DATE: 2000-02-24
 - 17 <160> NUMBER OF SEQ ID NOS: 49
 - 19 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

Does Not Comply
Corrected Diskette Needet

- 627 <210> SEQ ID NO: 49
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 632 <220> FEATURE:
- 633 <223> OTHER INFORMATION: Dyad symmetry sequence followed by a stretch of thymidine residue
 - 636 <400> SEQUENCE: 49
 - 637 gattctattt tttcttcgag tcacgctctc cccctaaagg agaaatgtga ctcaaaaaac 60
 - 639 aaaccttaaa aatcaaattt tagaaataag tcaaatttta gaaataagtc aaattttaga 120
 - 641 aataagtcaa attttagaaa taagtcaaat tttaataaag ctttatctta ttttaagaac 180
 - 643 tgtctacatt atagaccaca atgttattaa tttttagcta

220

E--> 646 016921-167.st25 E--> 651 Page 11

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/698,439A

DATE: 11/02/2004 TIME: 13:42:46

Input Set : A:\pto.lm.txt

Output Set: N:\CRF4\11022004\J698439A.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:646 M:254 E: No. of Bases conflict, LENGTH:Input:25 Counted:224 SEQ:49

L:646 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:11

L:646 M:112 C: (48) String data converted to lower case,

L:651 M:254 E: No. of Bases conflict, LENGTH:Input:11 Counted:228 SEQ:49 L:651 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2 L:651 M:252 E: No. of Seq. differs, <211> LENGTH:Input:220 Found:228 SEQ:49